

Result No.	Score	Query Match	Length	DB	ID	Description
1	4064.4	92.5	4126	15	US-10-241-220-17	Sequence 17, Appl
2	4029	91.7	4385	15	US-10-162-012-43	Sequence 43, Appl
3	4029	91.7	4385	15	US-10-144-624-1	Sequence 1, Appl
4	4029	91.7	4385	16	US-10-162-102-43	Sequence 43, Appl
5	3924.4	89.3	4075	15	US-10-328-198-1	Sequence 1, Appl
6	1871.8	42.6	1917	13	US-10-332-447-58	Sequence 58, Appl
7	1642.6	37.4	1714	16	US-10-120-988-437	Sequence 437, App
8	1625.2	37.0	1689	15	US-10-162-012-45	Sequence 45, Appl
9	1625.2	37.0	1689	15	US-10-144-624-3	Sequence 3, Appl
10	1625.2	37.0	1689	16	US-10-162-102-45	Sequence 45, Appl
11	1289	29.2	1284	15	US-10-029-386-20633	Sequence 20633, A
12	529.6	12.1	530	13	US-10-027-632-136458	Sequence 136458, S
13	529.6	12.1	530	16	US-10-027-632-136458	Sequence 136458, S
14	527.2	12.0	593	15	US-10-029-386-775	Sequence 775, App

QY 362 TTGACTTTGGGCTAAGCTGCTTGGAGCAGGAGTTCTTGGTGGGAGAGCTGCTCTCTGGGG 421
DB 121 TTGACTTTGGGCTAAGCTGCTTGGAGCAGGAGTTCTTGGTGGGAGAGCTGCTCTCTGGGG 180
QY 422 CTCTCTCGCTCCCTGCTTGGTGGCTTCTCATTTGACTGTATGGAGGAGCAAGCCA 481
DB 181 CTCTCTCGCTCCCTGCTTGGTGGCTTCTCATTTGACTGTATGGAGGAGCAAGCCA 240
QY 482 TCCTGGGAGCAACTTGGTGTCTGTGGAGGAGCCTGACCTGGGCTGCTGTGTTCCC 541
DB 241 TCCTGGGAGCAACTTGGTGTCTGTGGAGGAGCCTGACCTGGGCTGCTGTGTTCCC 300
QY 542 TGGCTGGCTGGCTCTGGGCGCGCTGTGGTGTGGCTTGGCACTTCCCTCTCTCCATGG 601
DB 301 TGGCTGGCTGGCTCTGGGCGCGCTGTGGTGTGGCTTGGCACTTCCCTCTCTCCATGG 360
QY 602 CTGTGTGTATCTAGGTGTAGAGCTGTGGGGCCACGGGAGCGGGAGTGTGTGTCCC 661
DB 361 CTGTGTGTATCTAGGTGTAGAGCTGTGGGGCCACGGGAGCGGGAGTGTGTGTCCC 420
QY 662 TCTATGAGGAGGAGCATCACCGTGGGATCTGTCTCTCTATGCGCTCAACTATGCACTGG 721
DB 421 TCTATGAGGAGGAGCATCACCGTGGGATCTGTCTCTCTATGCGCTCAACTATGCACTGG 480
QY 722 CTGTGACCCCTGGGATGGAGGACATGTTGGCTGGGCACTGCACTGTGTCTCTGTC 781
DB 481 CTGTGACCCCTGGGATGGAGGACATGTTGGCTGGGCACTGCACTGTGTCTCTGTC 540
QY 782 AATCCCTCAGCT 841
DB 541 AATCCCTCAGCT 600
QY 842 TCATCCCACTCCAGGAGGTGAGGCGCCCAAGCTGGGCGGGGAGGCGGAGTGTCTCTCT 901
DB 601 TCATCCCACTCCAGGAGGTGAGGCGCCCAAGCTGGGCGGGGAGGCGGAGTGTCTCTCT 660
QY 902 TTCTGAGACTCTTCAGGAGCGGATTAACATGCGAGGCGGAGGCGGAGTGTCTCTCTCT 961
DB 661 TTCTGAGACTCTTCAGGAGCGGATTAACATGCGAGGCGGAGGCGGAGTGTCTCTCTCT 720
QY 962 TGTGTCTCTTCAGCAACTAACAGGAGCAGCCCAAGCTGTGTGTGTGTGTGTGTGTGTGT 1021
DB 721 TGTGTCTCTTCAGCAACTAACAGGAGCAGCCCAAGCTGTGTGTGTGTGTGTGTGTGTGT 780
QY 1022 TCAGCTCCGTTGTTTCATGAGGAGTCTCAAGCTGTGTGTGTGTGTGTGTGTGTGTGTGT 1081
DB 781 TCAGCTCCGTTGTTTCATGAGGAGTCTCAAGCTGTGTGTGTGTGTGTGTGTGTGTGTGT 840
QY 1082 CAGTGAAGTGGCAGCTACCTCAGCGCCATGGGCTGTGTGTGTGTGTGTGTGTGTGTGTGT 1141
DB 841 CAGTGAAGTGGCAGCTACCTCAGCGCCATGGGCTGTGTGTGTGTGTGTGTGTGTGTGTGT 900
QY 1142 CTCTGTGTAGT 1201
DB 901 CTCTGTGTAGT 960
QY 1202 GCTTTGGCCCTGAGGAGT 1261
DB 961 GCTTTGGCCCTGAGGAGT 1020
QY 1262 AGACAGGCTCTCTGAGAGT 1321
DB 1021 AGACAGGCTCTCTGAGAGT 1080
QY 1322 GGACCAATGAGGAGCAAGGAGGAGCAATCTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGT 1381
DB 1081 GGACCAATGAGGAGCAAGGAGGAGCAATCTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGT 1140
QY 1382 CCAGATCTGGAGACCCCTCAGCCCT 1441
DB 1441 CCAGATCTGGAGACCCCTCAGCCCT 1200
QY 1442 CCCCTCTGCCGCTCGGGGGGATGCACTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGTGT 1501

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QY 1502 TCTTTGTGAGT 1561
DB 1261 TCTTTGTGAGT 1320
QY 1562 TCTTACCCTCTGAGATAGAGGAGGAGCTTGGCTTCTGTGCAACAGCTTCAACTGGGGGG 1621
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DB 1381 CCAACTCTTTCATCAGCCTGA 1440
QY 1682 CCTTCTCTCTCTACGGACTGACCGCTGTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 1741
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QY 1802 CCTGAGCTTGGGCAAGGAGGAGTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 1861
DB 1561 CCTGAGCTTGGGCAAGGAGGAGTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 1620
QY 1862 CTGGGCGCTCTGAGGAGTCT 1920
DB 1621 CTGGGCGCTCTGAGGAGTCT 1680
QY 1921 ATCTCTAGCT 1980
DB 1681 ATCTCTAGCT 1740
QY 1981 GGCCCTCTGCCCAAGGAGT 2040
DB 1741 GGCCCTCTGCCCAAGGAGT 1800
QY 2041 TGCCCAATCT 2100
DB 1801 TGCCCAATCT 1860
QY 2101 CTGGGTTTCCCAATGAGT 2160
DB 1861 CTGAGTTTCCCAATGAGT 1920
QY 2161 GTCTTTGTGCGCCATGAGT 2220
DB 1921 GTCTTTGTGCGCCATGAGT 1980
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DB 1981 GTCTTCTCCGATATACCCCTAAATCAATGAGGATATCATCTTTCTTAATCTCTTTTTT 2040
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DB 2101 TTTTGTGAGTGTGAGTGTCTCATTTGTGTGCGGAGGCTGTGTGTGTGTGTGTGTGTGTGTGT 2160
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DB 2161 CTCCACTTCTGTGTTCAAGCGATTTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 2220
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DB 2221 GGCGGTGCGCAACACCCAGCTAAATTTATTTTATGAGAGATGGGTTTCTACTGTGTGTG 2280
QY 2521 GCGAGGCTGTGTGAGT 2580

Db 2281 GCCAGGCTGGTGAATCTCTGAGCTCAAGTGATCCACCCACCTCAGCCTCCCAAGATG 2340
Qy 2581 CTAGGATTAACAGCCCTTTTGACTCTTTATCTGAGTTTTTATTTGAGCCCTCTAAATCTCTT 2640
Db 2341 CTAGGATTAACAGCCCTTTTGACTCTTTATCTGAGTTTTTATTTGAGCCCTCTAAATCTCTT 2400
Qy 2641 ACCCAGAATATTTATCTCTCACAGCAACTCTGACTCTTTTGAAGGAGGCTCAGTTCTA 2700
Db 2401 ACCCAGAATATTTATCTCTCACAGCAACTCTGACTCTTTGAGGAGGCTCAGTTCTA 2460
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Qy 3001 TAAATATGCTCCAGGGGTATATTTAGACCTGTGTTCTTTCCAGGAGGGTCCCAAGCTGT 3060
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Qy 3061 CCAGGGCTGGGAATTTCTACTATCTCTCATTTACCCAGGTCCTCTTTGGACCTGTA 3120
Db 2821 CCAGGGCTGGGAATTTCTACTATCTCTCATTTACCCAGGTCCTCTTTGGACCTGTA 2880
Qy 3121 AAGGGTCAGGGTGAATCAGATGGGGACTGAGCAAGTAGCTATGACTGCAGATCATGTAA 3180
Db 2881 AAGGGTCAGGGTGAATCAGATGGGGACTGAGCAAGTAGCTATGACTGCAGATCATGTAA 2940
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Db 3001 AGGTCTGATGCTTTGTTCATCCATGCTGTCATATGGTGTGGCAGAGCCCCCAGG 3060
Qy 3301 ACTCTGGCTCTCGAGTCTCTCTATCTCTCCATCTAGATGCTTCCCTTGTATCCAGTG 3360
Db 3061 ACTCTGGCTCTCGAGTCTCTCTATCTCTCCATCTAGATGCTTCCCTTGTATCCAGTG 3120
Qy 3361 ATGTCTGGAGCTGGCTTTGCCAAGCTGTGTGAGAGCTGTTGCTACATTTTCAGGATTTT 3420
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Db 3241 AATTACATTTAAACAAAAAATTAATCTCAAAATTCATTTACTTAAATTTTACTACCTGTTA 3300
Qy 3541 CTATTATCTGCTTTTGGGCTATTTCTACATAGTAACCTTTATGAGACCTTAGGGAG 3600
Db 3301 CTATTATCTGCTTTTGGGCTATTTCTACATAGTAACCTTTATGAGACCTTAGGGAG 3360
Qy 3601 ACACCGGCTATCTCTCTGATCCCACTCAATGACATCATGCTTAGTCTTTGGTGTCTT 3660
Db 3361 ACACCGGCTATCTCTCTGATCCCACTCAATGACATCATGCTTAGTCTTTGGTGTCTT 3420

Qy 3661 AACTGGCTGTGGGAGTGTGTTTTGTATCAAAAAGATTAGAGAGGACTACATCATCAGGCT 3720
Db 3421 AACTGGCTGTGGGAGTGTGTTTTGTATCAAAAAGATTAGAGAGGACTACATCATCAGGCT 3480
Qy 3721 TGATTTATGTTGTTGATTTTTCTAGACTTCAGAACATGCTGGATAAAATGTCAGTAATG 3780
Db 3481 TGATTTATGTTGTTGATTTTTCTAGACTTCAGAACATGCTGGATAAAATGTCAGTAATG 3540
Qy 3781 CAATTTAACTTTAAAGTATGTTCTGTTGTTAGCCAAATACATGGTGTATAGCACCAAAA 3840
Db 3541 CAATTTAACTTTAAAGTATGTTCTGTTGTTAGCCAAATACATGGTGTATAGCACCAAAA 3600
Qy 3841 ATGAGGGATTAATCTCTCCAGTAGTTGAACACTGTTCATCCGTTTCAGCTGACAGCTGCTC 3900
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Qy 3901 AATCATTTAAGAGGAGTTCTGACATTTTCATTTGTTTACTTTGTTCTCTCTCNC 3960
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Qy 4021 TAGCTGTACAGTTATCAGGGATTTTATTTCTGAGTCTAATTTTGTCAAATCATTGGCCAAA 4080
Db 3781 TAGCTGTACAGTTATCAGGGATTTTATTTTGTAGTCTAATTTTGTCAAATCATTGGCCAAA 3840
Qy 4081 TCGCAGTATAGTTGACTTTTGAGTACAAGGTTTGGCAAAAAAATAATTAACAATA 4140
Db 3841 TCGCAGTATAGTTGACTTTTGAGTACAAGGTTTGGCAAAAAAATAATTAACAATA 3900
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Db 3901 TTCTGTAAGAATCAATTCGCTATATGGAATTTAGGATAAAGAAATTTTACAAATAAGAAAT 3960
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Db 3961 ATTTACAATAAGAGTATTTATTTATTTTGAAGTTGTGCAACAAACATACCCCTTTATC 4020
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Qy 4321 TGGAAATTTAGGATAGAATATTTACAAATAAGAGTATTTTACAAATAA 4366
Db 4081 TGGAAATTTAGGATAGAATATTTTACAAATAAGAGTATTTTACAAATAA 4126

RESULT 2

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; GENERAL INFORMATION:

; APPLICANT: Curtis, Rory A.J.

; APPLICANT: Silos-Santiago, Inmaculada

; APPLICANT: Gu, Wei

; TITLE OF INVENTION: NOVEL HUMAN ION CHANNEL AND TRANSPORTER FAMILY MEMBERS

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; PRIOR FILING DATE: 2000-06-05

; PRIOR APPLICATION NUMBER: US 09/875,423

; PRIOR FILING DATE: 2001-06-05

; PRIOR APPLICATION NUMBER: PCT/US01/18398

Db	1725	GTTCCTGAAACAAAGGCCAGTGGTTGGCAGAGATAGACCGACGAGTGTCCAGACAGACGCG	1784
Qy	1798	TTCAACCTCGAGCTTTTGGCCACAGGCAGAACTCCACTGGCATCCCGTACAGCCGCACTCGAG	1857
Db	1785	TTCAACCTCGAGCTTTTGGCCACAGGCAGAACTCCACTGGCATCCCGTACAGCCGCACTCGAG	1844
Qy	1858	ATCTCTGGGGCTCTCTGAGGAAATCCGTCTGGCTGG--AATTCTGGAACTGTGGCTTTGGCA	1916
Db	1845	ATCTCTGGGGCTCTCTGAGGAAATCCGTCTGGCTGGAAATTTCTGAACTGTGGCTTTGGCA	1904
Qy	1917	GACCATCTTCAGCATCTGCTTCTAGCCCCAGAGCACAAAGTTTCCAGCTGCTCTTTTGG	1976
Db	1905	GACCATCTTCAGCATCTGCTTCTAGCCCCAGAGCACAAAGTTTCCAGCTGCTCTTTTGG	1964
Qy	1977	GAGTGGCCCTCGCCCCAAAGTGGTTTGTCTTTTGTGGGTAAAAAGGATGAAGATTTTG	2036
Db	1965	GAGTGGCCCTCGCCCCAAAGTGGTCTGCTTTTGTCTGGGTAAAAAGGATGAAGTCTG	2024
Qy	2037	AGAAATGCCCAATTTCTCAATTTTGGGTTCAGGCCCTGAAGGTTCTTGAGGATCTAGTTTC	2096
Db	2025	AGAAATGCCCAACTCTTCAATTTTGAAGTCTCAGGCCCTGAAGGTTCTTGAGGATCTAGCTTC	2084
Qy	2097	ATGCGCTCGGTTTCCCATAGACTTGGACATTTTGTGACATTTTGTAGATTTTATAGAGAAATATCTA	2156
Db	2085	ATGCGCTCAGTTTCCCATAGACTTGGACATCTCTGACATATTTTATAGAGAAATATTTCTA	2144
Qy	2157	TGAAGTCTTTTGTGCCCATAGATTTTCTTTTCAAAGAATCTCAGGGGTACCAATCCGGGCA	2216
Db	2145	TGAAGTCTTTTGTGCCCATAGGACTTTTCTCAAGAATCTCAGGGGTACCAATCTGGCA	2204
Qy	2217	GGAGGTTTTTCCGATATCACCCCTAAATCCAAATGAGGATATCATCTTTTCTTAATCTCT	2276
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Db	2383	CAACCTCCACTCTCTGAGTTCAAGCGAATCTTGTGCTCGCCTCTTAAGCAGCTGGAC	2442
Qy	2457	TACAGGGCGGTGCCACACACCCAGCTAAATTTATTTTTTACGAGAGATGGGGTTTCACTGT	2516
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Qy	2577	AGTCTAGGAATACAGGCCCTTTTGACTCTTTTTATCTGAGTTTATTGACCCCTCTAATTC	2636
Db	2563	AGTCTAGGAATACAGGCCCTTTTGACTCTTTTTATCTGAGTTTATTGACCCCTCTAATTC	2622
Qy	2637	TCATTACCCAGAAATATTTATCTTCACAGCAATCTGACTCTTTTGACGGGAGGCTCAGT	2696
Db	2623	TCATTACCCAGAAATATTTATCCCTTACCAGCAACTCTGACTCTTTTGACGGGAGGCTCAGT	2682
Qy	2697	CTAGTCTCTGGTCTGTGGTGTCATTTGCTGTAGGAATGACCAAGGCTCAGTTTCCCC	2756
Db	2683	CTAGTCTCTGGTCTGTGGTGTCATTTGCTGTAGGAATGACCAAGGCTCAGTTTCCCC	2742
Qy	2757	ATTTGTATATGGGAAGCCTGTACAGGCTCAATCTTAAGATTTTCTCTGACTCCAGTGAG	2816
Db	2743	ATTTGTATATGGGAAGCCTGTACAGGCTCAATCTTAAGATTTTCTCTGACTCCAGTGAG	2802
Qy	2817	CTGAAATTTCAAATGCTGGTCTAGGAGCTGTCTCCAGGATGGTGCAAGATGGCTTTGCGG	2876
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Qy	2877	AAAGGAGATGGGTTTGGAGGCCAACAAACCTGCTGTGTCAAATATTTGGCCTTTGGCCTTTGGC	2936
Db	2863	AAAGGAGATGGGTTTGGAGGCCAACAAACCTGCTGTGTCAAATATTTGGCCTTTGGCCTTTGGC	2922
Qy	2937	AGCCCTTGAACCTTGAGTAABATACAACTCCCTGAACTCAGTTTCCCTCATCTCGAGATG	2996
Db	2923	AGCCCTTGAACCTTGAGTAABATACAACTCCCTGAACTCAGTTTCCCTCATCTCGAGATG	2982
Qy	2997	GGGATAATATATGCCCCAGGGGTATATTTAGACCCCTGTTTCCCTTTCAGAGAGGGTCCCCAGC	3056
Db	2983	GGGATAATATATGCCCCAGGGGTATATTTAGACCCCTGTTTCCCTTTCAGAGAGGGTCCCCAGC	3042
Qy	3057	TGCTCAGGGCTGGGAAATTTCTATTATCTCTCATTTACCCAGGTCCTCTCTTTTGAACCC	3116
Db	3043	TGCTCAGGGCTGGGAAATTTCTATTATCTCTCATTTACCCAGGTCCTCTCTTTTGAACCC	3102
Qy	3117	TGTAAGGGGTCAAGGTCAATCAGATGGGAGCTGAGCAAGTAGCTATGACTGCAGATCAT	3176
Db	3103	TGTAAGGGGTCAAGGTGNAATCAGATGGGAGCTGAGCAAGTAGCTATGACCGCAGATCAT	3162
Qy	3177	GTAAGGAAGGAGCTGACAAGAAGCTCCAGATGCTGGGAGAAATGAAGAGCTAAAATAGA	3236
Db	3163	GTAAGGAAGGAGCTGACAAGAAGCTCCAGATGCTGGGAGAAATGAAGAGCTAAAATAGA	3222
Qy	3237	TCCTAGGTGCTGATGCTTTTGTCAATCCATCGGTGCACATATGGTGTCTGCAGAGCCCCC	3296
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Qy	3297	AAGGACCTCGCCCTCTCGAGTTCTCTCTATCTTCTCCATCTCTAGATGCTCCCTTTGATACC	3356
Db	3283	AAGGACCTCTGGCCTCTCGAGTTCTCTCTATCTTCTCCATCTCTAGATGCTTCCCTTTGATACC	3342
Qy	3357	AGTGATGCTGGAGCTGGCTTTGCCAAGCTGTGAGAGCTGTGTCTACATTTTTCAGGA	3416
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Qy	3417	TTTTTACAAGTGTGTAAACACAGCCATTATAAAAAATTAATGATTTTAAATTTTATAATTA	3476
Db	3403	TTTTTACAAGTGTGTAAACACAGCCATTATAAAAAATTAATGATTTTAAATTTTATAATTA	3462
Qy	3477	AGTAAATTTACATTTAAAAACAAAAATTTATACTCATAATTCATTTACTTTTACTACCT	3536
Db	3463	AGTAAATTTACATTTAAAAACAAAAATTTATACTCATAATTCATTTACTTTTACTACCT	3522
Qy	3537	GTTACTATTATCTGTGCTTTTGAGGCTATTTTCTACATAGTAACTCTTTATGGAGACCTAGG	3596
Db	3523	GTTACTATTATCTGTGCTTTTGAGGCTATTTTCTACATAGTAACTCTTTATGGAGACCTAGG	3582
Qy	3597	GGAGACACGGGATCTCTTCTGATTTCCCCACCTCAATGACATCATGTGTAGTCTTTGGTT	3656
Db	3583	GGAGACACGGGATCTCTTCTGATTTCCCCACCTCAATGACATCATGTGTAGTCTTTGGTT	3642
Qy	3657	GCTTTAACTGGCTGTGGGAGTGTTTTTTGATATCAAAAGATTTAGAGAGGACTACACATCAG	3716
Db	3643	GCTTTAACTGGCTGTGGGAGTGTTTTTTGATATCAAAAGATTTAGAGAGGACTACACATCAG	3702
Qy	3717	GGCTTGATTTATTTGTTTGTGATTTTCTAGACTTCAGAAATGCTGTGATTAATGTCTAGT	3776
Db	3703	GGCTTGATTTATTTGTTTGTGATTTTCTAGACTTCAGAAATGCTGTGATTAATGTCTAGT	3762
Qy	3777	AATGCAAAATTAACCTTTTAAAGTATGCTCTGTTTGTAGCCAAATACATGGTGTATAGCACCA	3836
Db	3763	AATGCAAAATTAACCTTTTAAAGTATGCTCTGTTTGTAGCCAAATACATGGTGTATAGCACCA	3822
Qy	3837	AAAAATGGAGGATTTATTTCTCCAGTAGTTTGAACACTGTCTCCGTTTTCAGCTGCACAGCT	3896
Db	3823	AAAAATGGAGGATTTATTTCTCCAGTAGTTTGAACACTGTCTCCGTTTTCAGCTGCACAGCT	3882
Qy	3897	GCTCAAAATCATTTAAGAAGAGTCTGCACATTTTCAATTTTCAATTTTGTCTTTTGTCTTCC	3956
Db	3883	GCTCAAAATCATTTAAGAAGAGTCTGCACATTTTCAATTTTCAATTTTGTCTTTTGTCTTCC	3942

1078 GGCGCAGTCAAGGTGGCAGCTACCCCTGACCGCCATGGGGCTGGTGAACGCTGCGAGGCCG 1137
Db |||||
1065 GGCGCAGTCAAGGTGGCAGCTACCCCTGACCGCCATGGGGCTGGTGAACGCTGCGAGGCCG 1124
Qy |||||
1138 AGGGCTCTTTGCTAGCTGGCTGTGCTCTCATGGCCCTGTCCGTCACTGAGTGGCATAGGCCCTC 1197
Db |||||
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Qy |||||
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Db |||||
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Qy |||||
1378 CATCCAGACTGGAGACCCCTCAGCCCTCTCGCTGGCCCTGAGCTCTGCCCTCCCT 1437
Db |||||
1365 CATCCAGACTGGAGACCCCTCAGCCCTCTCGCTGGCCCTGAGCTCTGCCCTCCCT 1424
Qy |||||
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1965 GAGTGGCCCTCTGCCCCCAAGGTGGTTTCTTTTCTGGGGTAAAGAGGATGAAAGTCTG 2024
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RESULT 6
US-10-332-447-58
; Sequence 58, Application US/10332447
; Publication No. US20040053258A1
; GENERAL INFORMATION:
; APPLICANT: INCYTE GENOMICS, INC.; RAUMANN, Brigitte E.;
; APPLICANT: THORNTON, Michael; DING, Li; YUE, Henry;
; APPLICANT: TANG, Y.Tom; HARLAND, Lee; BURFORD, Neil;
; APPLICANT: GREENE, Barrie D.; SANJANWALA, Madhu S.;
; APPLICANT: BAUGHN, Mariah R.; YAO, Monique G.; YANG, Junming;
; APPLICANT: ARVIZU, Chandra S.; GANDHI, Ameena R.;
; APPLICANT: HAFALIA, April J.A.; TRIBOULEY, Catherine M.;
; APPLICANT: WALIA, Nandier K.; AU-YOUNG, Janice;
; APPLICANT: WALSH, Roderick T.; RAMKUMAR, Jayalaxmi;
; APPLICANT: LU, Yan; LU, Dyung Aina M.; AZIMZAI, Valda;
; APPLICANT: LU, Yuming; SEILHAMER, Jeffrey J.; BOROWSKI, Mark L.;
; APPLICANT: KHAN, Farrah A.; KEARNEY, Liam; THANGAVELU, Kavitha;
; APPLICANT: DAS, Debopriya; POLICKY, Jennifer L.
; TITLE OF INVENTION: TRANSPORTERS AND ION CHANNELS
; FILE REFERENCE: PI-0149 USN
; CURRENT APPLICATION NUMBER: US/10/332,447
; CURRENT FILING DATE: 2003-01-07
; PRIOR APPLICATION NUMBER: US 60/216,547
; PRIOR FILING DATE: 2000-07-07
; PRIOR APPLICATION NUMBER: US 60/218,232
; PRIOR FILING DATE: 2000-07-14
; PRIOR APPLICATION NUMBER: US 60/220,112
; PRIOR FILING DATE: 2000-07-21
; PRIOR APPLICATION NUMBER: US 60/221,839
; PRIOR FILING DATE: 2000-07-28
; NUMBER OF SEQ ID NOS: 64
; SOFTWARE: PERL Program
; SEQ ID NO 58
; LENGTH: 1917
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc feature
; OTHER INFORMATION: Incyte ID No. US20040053258A1 5047435CB1
US-10-332-447-58
Query Match 42.6%; Score 1871.8; DB 13; Length 1917;
Best Local Similarity 99.6%; Pred. No. 0;
Matches 1887; Conservative 0; Mismatches 7; Indels 1; Gaps 1;
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FILE REFERENCE: 108627.129
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-04-20
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR FILING DATE: 1999-11-23
PRIOR APPLICATION NUMBER: US 60/156,358
PRIOR FILING DATE: 1999-09-28
PRIOR APPLICATION NUMBER: US 60/146,002
PRIOR FILING DATE: 1999-08-09
NUMBER OF SEQ ID NOS: 325720
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 136458
LENGTH: 530
TYPE: DNA
ORGANISM: Human
US-10-027-632-136458

Query Match 12.1%; Score 529.6; DB 13; Length 530;
Best Local Similarity 99.8%; Pred. No. 1.6e-113;
Matches 529; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

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QY 3178 TAAAGGAGGGACTGACAAGAGCTCCAGATGCTGGGAGAAATGAAGAGCTAAATATAGAT 3237
DB 121 TAAAGGAGGGACTGACAAGAGCTCCAGATGCTGGGAGAAATGAAGAGCTAAATATAGAT 180

QY 3238 CCTAGGTGCTGATGCTTTGTTCATCGCTGCACATATGGTGTGTCAGAGCCCA 3297
DB 181 CCTAGGTGCTGATGCTTTGTTCATCGCTGCACATATGGTGTGTCAGAGCCCA 240

QY 3298 AGGACTCTGGCCTCTCGAGTCTCTCTATCTTCCATCTAGATGCTTCCCTTTGATCCA 3357
DB 241 AGGACTCTGGCCTCTCGAGTCTCTCTATCTTCCATCTAGATGCTTCCCTTTGATCCA 300

QY 3358 GTGATGCTGAGCTGGCTTTGCCAAGCTTTGTGAGAGCTGGTGTGCTACATTTTCAGGAT 3417
DB 301 GTGATGCTGAGCTGGCTTTGCCAAGCTTTGTGAGAGCTGGTGTGCTACATTTTCAGGAT 360

QY 3418 TTTTACAAGTTGGTAAACACAGCCATTATAAAAAATTAATGATTTAAATTTAATTTAA 3477
DB 361 TTTTACAAGTTGGTAAACACAGCCATTATAAAAAATTAATGATTTAAATTTAATTTAA 420

QY 3478 GTAAATTTACATTTAAACAAAAAATTTATCTCAAAATTCATTACTTTAATTTTACTACCTG 3537
DB 421 GTAAATTTACATTTAAACAAAAAATTTATCTCAAAATTCATTACTTTAATTTTACTACCTG 480

QY 3538 TTACTATTATCTGCTTTTGGGCTATTTTCTACATAGTAACCTTTATGG 3587
DB 481 TTACTATTATCTGCTTTTGGGCTATTTTCTACATAGTAACCTTTATGG 530
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RESULT 13

US-10-027-632-136458
Sequence 136458, Application US/10027632
Publication No. US20030204075A9
GENERAL INFORMATION:
APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
Polymorphisms in the Human Genome

RESULT 14

US-10-029-386-775/c
Sequence 775, Application US/10029386
Publication No. US20030194704A1
GENERAL INFORMATION:
APPLICANT: Penn, Sharron G.
APPLICANT: Rank, David R.

FILE REFERENCE: 108627.129
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-04-20
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR FILING DATE: 1999-11-23
PRIOR APPLICATION NUMBER: US 60/156,358
PRIOR FILING DATE: 1999-09-28
PRIOR APPLICATION NUMBER: US 60/146,002
PRIOR FILING DATE: 1999-08-09
NUMBER OF SEQ ID NOS: 325720
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 136458
LENGTH: 530
TYPE: DNA
ORGANISM: Human
US-10-027-632-136458

Query Match 12.1%; Score 529.6; DB 16; Length 530;
Best Local Similarity 99.8%; Pred. No. 1.6e-113;
Matches 529; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

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QY 3058 GGTCCAGGGCCTGGGAAATTTCTACTTATCCTCATTTACCCAGGTCCTCTTTGGACCT 3117
DB 1 GGTCCAGGGCCTGGGAAATTTCTACTTATCCTCATTTACCCAGGTCCTCTTTGGACCT 60

QY 3118 GTAAAGGGTCAGGGTGAATCAGATGGGGACTGAGCAAGTACTGATGTCAGATCATG 3177
DB 61 GTAAAGGGTCAGGGTGAATCAGATGGGGACTGAGCAAGTACTGATGTCAGATCATG 120

QY 3178 TAAAGGAGGGACTGACAAGAGCTCCAGATGCTGGGAGAAATGAAGAGCTAAATATAGAT 3237
DB 121 TAAAGGAGGGACTGACAAGAGCTCCAGATGCTGGGAGAAATGAAGAGCTAAATATAGAT 180

QY 3238 CCTAGGTGCTGATGCTTTGTTCATCGCTGCACATATGGTGTGTCAGAGCCCA 3297
DB 181 CCTAGGTGCTGATGCTTTGTTCATCGCTGCACATATGGTGTGTCAGAGCCCA 240

QY 3298 AGGACTCTGGCCTCTCGAGTCTCTCTATCTTCCATCTAGATGCTTCCCTTTGATCCA 3357
DB 241 AGGACTCTGGCCTCTCGAGTCTCTCTATCTTCCATCTAGATGCTTCCCTTTGATCCA 300

QY 3358 GTGATGCTGAGCTGGCTTTGCCAAGCTTTGTGAGAGCTGGTGTGCTACATTTTCAGGAT 3417
DB 301 GTGATGCTGAGCTGGCTTTGCCAAGCTTTGTGAGAGCTGGTGTGCTACATTTTCAGGAT 360

QY 3418 TTTTACAAGTTGGTAAACACAGCCATTATAAAAAATTAATGATTTAAATTTAATTTAA 3477
DB 361 TTTTACAAGTTGGTAAACACAGCCATTATAAAAAATTAATGATTTAAATTTAATTTAA 420

QY 3478 GTAAATTTACATTTAAACAAAAAATTTATCTCAAAATTCATTACTTTAATTTTACTACCTG 3537
DB 421 GTAAATTTACATTTAAACAAAAAATTTATCTCAAAATTCATTACTTTAATTTTACTACCTG 480

QY 3538 TTACTATTATCTGCTTTTGGGCTATTTTCTACATAGTAACCTTTATGG 3587
DB 481 TTACTATTATCTGCTTTTGGGCTATTTTCTACATAGTAACCTTTATGG 530
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RESULT 14

US-10-029-386-775/c
Sequence 775, Application US/10029386
Publication No. US20030194704A1
GENERAL INFORMATION:
APPLICANT: Penn, Sharron G.
APPLICANT: Rank, David R.

APPLICANT: Hanzel, David K.
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
FILE REFERENCE: AOMICA-X-2
CURRENT APPLICATION NUMBER: US/10/029,386
CURRENT FILING DATE: 2001-12-20
NUMBER OF SEQ ID NOS: 34288
SOFTWARE: Annonax Sequence Listing Engine vers. 1.1
SEQ ID NO 775
LENGTH: 593
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
OTHER INFORMATION: MAP TO ALL33520.2
OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 2.7
OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 1.5
OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 3.1
OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 2.5
OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 1.9
OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 1.7
OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 1.7
OTHER INFORMATION: SWISSPROT HIT: P27546, EVALUE 3.00e-01
OTHER INFORMATION: EST HUMAN HIT: B1824797.1, EVALUE 0.00e+00
OTHER INFORMATION: NT HIT: G114786183, EVALUE 0.00e+00
US-10-029-386-775

Query Match 12.0%; Score 527.2; DB 15; Length 593;
Best Local Similarity 98.5%; Pred. No. 6.3e-113;
Matches 532; Conservative 0; Mismatches 8; Indels 0; Gaps 0;
QY 1014 CACATCTTCAGCTCCGTTGGTTCCATGGGGGATCCTCAGCGGTCTGGCTCTGTGGG 1073
DB 593 CACATCTTCAGCTCCGTTGGTTCCATGGGGGATCCTCAGCGGTCTGGCTCTGTGGG 534
QY 1074 GCTTGGCGGAGTCAAGTGCACCTACCTGACCGGCATGGGTGGTGGACGGTGCAGG 1133
DB 533 GCTTGGCGGAGTCAAGTGCACCTACCTGACCGGCATGGGTGGTGGACGGTGCAGG 474
QY 1134 CCGCAGGGCTCTGTTGCTAGCTGGCTGTGCTCTCATGGCCCTGTGCTCGTCAAGTGGCATAGG 1193
DB 473 CCGCAGGGCTCTGTTGCTAGCTGGCTGTGCTCTCATGGCCCTGTGCTCGTCAAGTGGCATAGG 414
QY 1194 CTTGCTCAGCTTTCGCTGCGCCATGACCTCAGCGCCCAAGCTGTGCTGCTGCGCCCAATGC 1253
DB 413 CTTGCTCAGCTTTCGCTGCGCCATGACCTCAGCGCCCAAGCTGTGCTGCTGCGCCCAATGC 354
QY 1254 CACCGGGCAGACAGGCTCTCCTGGAGACTCTGCGCTGCTGCGAGGACTCTCTACCTCC 1313
DB 353 CACCGGGCAGACAGGCTCTCCTGGAGACTCTGCGCTGCTGCGAGGACTCTCTACCTCC 294
QY 1314 CATTCAGGACCAATGAGGACCAAGGGAGGCAATCTTGTCCACTGTGTAAGAAACCAA 1373
DB 293 CATTCAGGACCAATGAGGACCAAGGGAGGCAATCTTGTCCACTGTGTAAGAAACCAA 234
QY 1374 GCCCATCCAGATCTGGAGACCCCTCAGCCCTCTGCGCTGCGAGGACTCTGCTGCTGCTGCTG 1433
DB 233 GCCCATCCAGATCTGGAGACCCCTCAGCCCTCTGCGCTGCGAGGACTCTGCTGCTGCTGCTG 174
QY 1434 CCTGGGCCCCCTCTGCGCCCTCGGGGGCATGCACTGCTGCGCTGCGAGGCGCACTGCTGTG 1493
DB 173 CCTGGGCCCCCTCTGCGCCCTCGGGGGCATGCACTGCTGCGCTGCGAGGCGCACTGCTGTG 114
QY 1494 CTTGATGCTTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 1553
DB 113 CTTGATGCTTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 54

RESULT 15
US-09-880-107-850/c
Sequence 850, Application US/09880107
Patent No. US20020142981A1
GENERAL INFORMATION:
APPLICANT: Horne, Darci T.

APPLICANT: Vockley, Joseph G.
APPLICANT: Scherf, Uwe
APPLICANT: Gene Logic, Inc.
TITLE OF INVENTION: Gene Expression Profiles in Liver Cancer
FILE REFERENCE: 44921-5028-WO
CURRENT APPLICATION NUMBER: US/09/880,107
CURRENT FILING DATE: 2001-06-14
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR FILING DATE: 2000-06-14
PRIOR APPLICATION NUMBER: US 60/237,054
PRIOR FILING DATE: 2000-10-02
NUMBER OF SEQ ID NOS: 3950
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 850
LENGTH: 528
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
OTHER INFORMATION: Genbank Accession No. US20020142981A1 AA404352
US-09-880-107-850
Query Match 12.0%; Score 526.4; DB 9; Length 528;
Best Local Similarity 99.8%; Pred. No. 8.9e-113;
Matches 527; Conservative 0; Mismatches 1; Indels 0; Gaps 0;
QY 3865 TTGAACACTGTCATCCGTTTCAGCTGACAGCTGCTCAATCATTTAAGAGGAGTCTTGA 3924
DB 528 TTGAACACTGTCATCCGTTTCAGCTGACAGCTGCTCAATCATTTAAGAGGAGTCTTGA 469
QY 3925 CATTCAATTTTCATTTGTTTACTTTTCTCTCACTAGTGTAAACAAAAATTTCAACCA 3984
DB 468 CATTCAATTTTCATTTGTTTACTTTTCTCTCACTAGTGTAAACAAAAATTTCAACCA 409
QY 3985 GCATTCATGCGCAACCTATACCCATTTCTCAGTGCCTAGCTGTACAGTATCAGGGATT 4044
DB 408 GCATTCATGCGCAACCTATACCCATTTCTCAGTGCCTAGCTGTACAGTATCAGGGATT 349
QY 4045 TTATTCGTAGTCTAAATTTTGTCAATCATGCGCCAAATCGCAGTGTAGTGTGACTTTGGAT 4104
DB 348 TTATTCGTAGTCTAAATTTTGTCAATCATGCGCCAAATCGCAGTGTAGTGTGACTTTGGAT 289
QY 4105 ACAAGGTTTGGCAAAAAAATAATTTAACAATAATTTCTGTAAAGATCAATTTGGCTATA 4164
DB 288 ACAAGGTTTGGCAAAAAAATAATTTAACAATAATTTCTGTAAAGATCAATTTGGCTATA 229
QY 4165 TGGAAATTTAGGATAAAGAAATTTTACAAATAAGAAATTTTACAAATAAGAAATTTTATTT 4224
DB 228 TGGAAATTTAGGATAAAGAAATTTTACAAATAAGAAATTTTACAAATAAGAAATTTTATTT 169
QY 4225 ATTTGTAAGTTGTGCAACAACAATACCCCTTTATCTCTGTAAATTTTATACACAAAA 4284
DB 168 ATTTGTAAGTTGTGCAACAACAATACCCCTTTATCTCTGTAAATTTTATACACAAAA 109
QY 4285 ATTTAACAAGATTTCTGTAAGAAATTTAATTTGGCTATATGGAATTTAGGATAGATTTTAC 4344
DB 108 ATTTAACAAGATTTCTGTAAGAAATTTAATTTGGCTATATGGAATTTAGGATAGATTTTAC 49
QY 4345 AATAAAGATTTTACAAATAAGGTTTGTATTTATTTTGTAAAAAAA 4392
DB 48 AATAAAGATTTTACAAATAAGGTTTGTATTTATTTTGTAAAAAAA 1

Search completed: June 23, 2004, 21:41:17
Job time : 1711 secs